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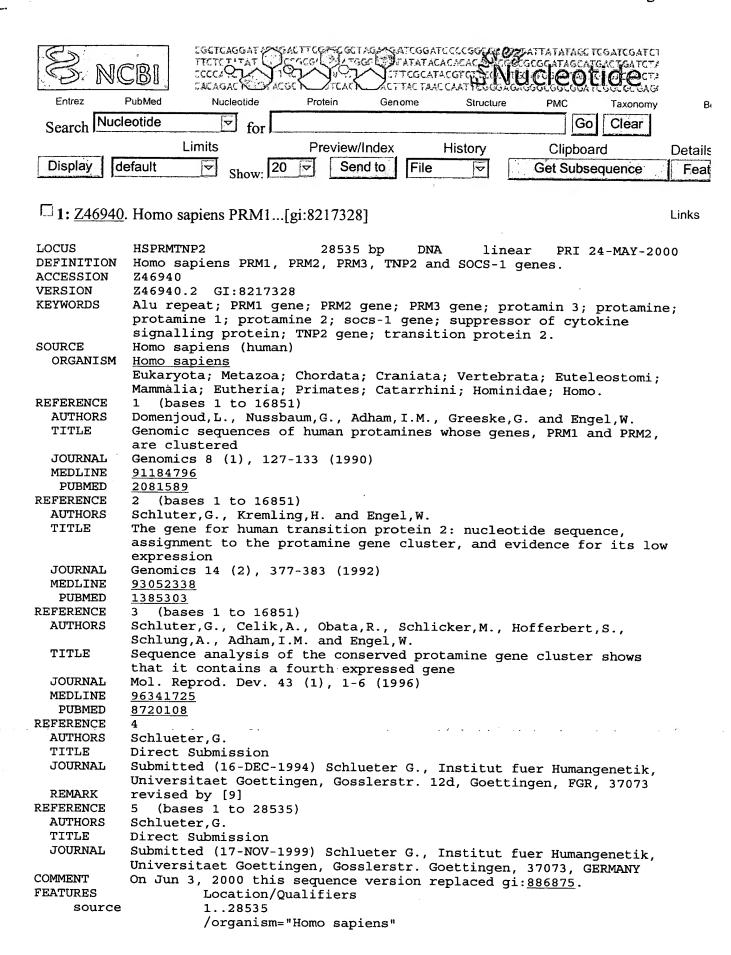
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HSPRWINP2 28535 bp DNA PRI 24-MAY-2000 Homo sapiens PRM1, PRM2, PRM3, TNP2 and SOCS-1 genes. 246940	246940.7 GI:8217328 Alu repeat; PRM1 gene; PRM2 gene; PRM3 gene; protamin 3; protamine; protamine 1; protamine 2; socs-1 gene; suppressor of cytokine	Signalling Protein; TNP2 gene; transition protein 2. human Homo sapiens Eukaryota; Metazoa; Chordata; Craniata: Vertebrata: Euteleostomi	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 16851) Domenjoud, L., Nussbaum, G., Adham, I.M., Greeske, G. and Engel, W. Genomic sequences of human protramines whom a new and and an and a new and a n	are clustered Genomics 8 (1), 127-133 (1990)	2 (bases 1 to 16851) Schluter, G., Kremling, H. and Engel, W. The gene for human transition protein 2: mucleatide accuracy	assignment to the protamine gene cluster, and evidence for its low, expression Genomics 14 (2), 377-383 (1992) 93052338
LOCUS DEFINITION ACCESSION	VERSION KEYWORDS	SOURCE ORGANISM	REFERENCE AUTHORS TITLE	JOURNAL	REFERENCE AUTHORS TITLE	JOURNAL
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27586

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gccgctttcacctggatggcagccgcgagagcttcgactgcctcttcgagctgctggagc

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27946 GATGCCGTGAAGCCGGTGCTGGGAAGTAGTTGATGTGCCGTTGCAGATTGGG

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						ttttttaaaaaaa 1094 	AB000734 1216 bp Homo sapièns mRNA for AB000734 AB000734.1 GI:2627028	
y 721 b 28066	y 781 b 28126	y 841 b 28186	Oy 901	y 961 b 28306	y 1021 b 28366	y 1081 b 28426	RESULT 10 ABO00734 LOCUS DEFINITION ACCESSION VERSION	KEYWORDS SOURCE ORGANISM
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Length 28535; Indels

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Query Match
Best Local Similarity 99.5
Matches 1089; Conservative

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